

Figure 1

1 RTPSDKPV AH VVANPQAE GQ LQWLNRRANA LLANGVELRD NQLVVPSEGL
51 YLIYSQVL FK GQGCPSTHVL LTHTISR IAV SYQTKVNLLS AIKSPCQRET
101 PEGAEAKPWY EPIYLGGV FQ LEKGDRLSAE INRPDYLLFA ESGQVYFGII
151 AL

Figure 2

1 HSDPARRGQL SVCDSISEWV TAADKKTAVD MSGGTVTVLE KVPVSKGQLK
51 QYFYETKCNP MGYTKEGCRG IDKRHWNSQC RTTQSYVRAL TMDSKKRIGW
101 RFIRIDTSCV CTLTIKRGR

FIGURE 3

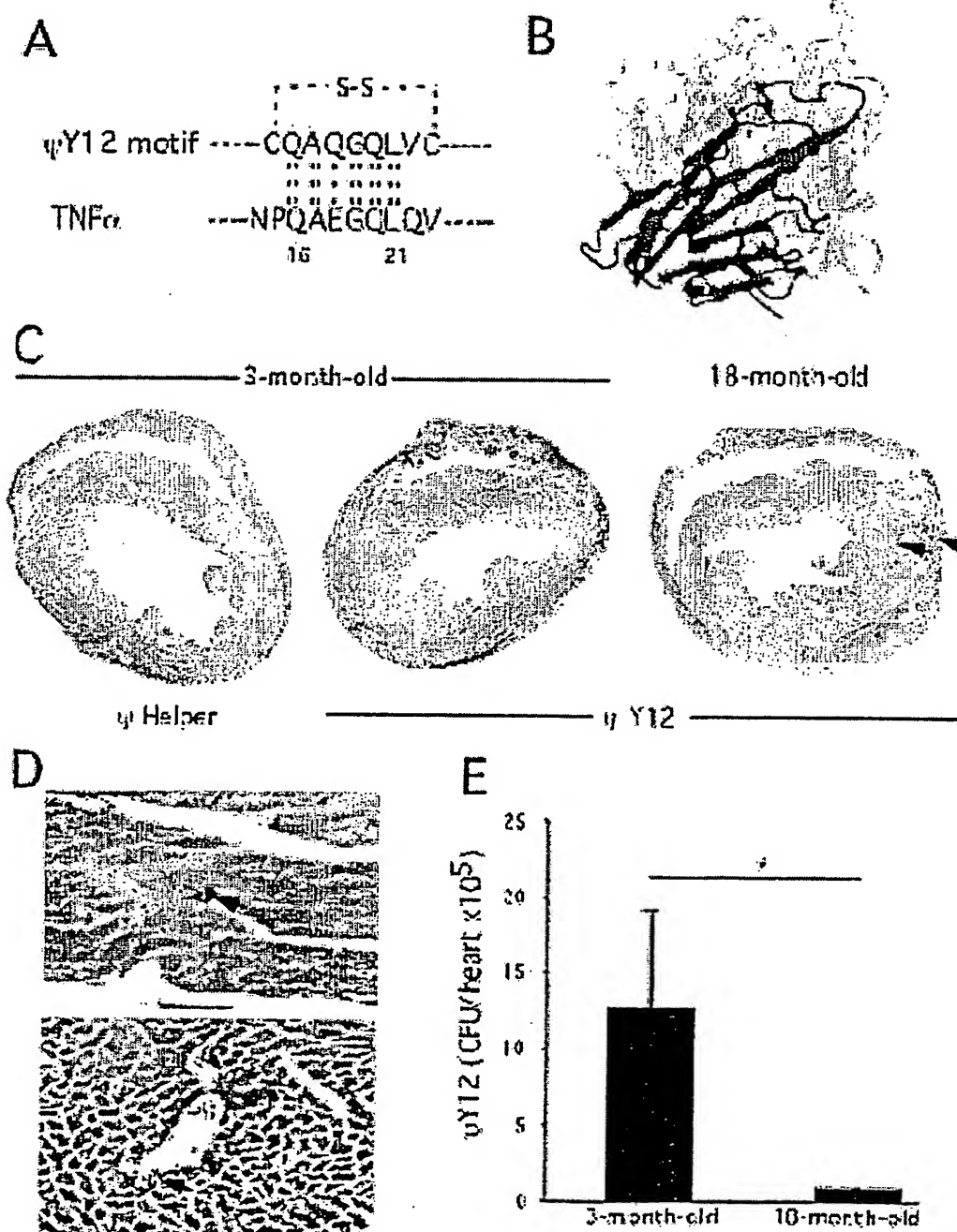


FIGURE 4A

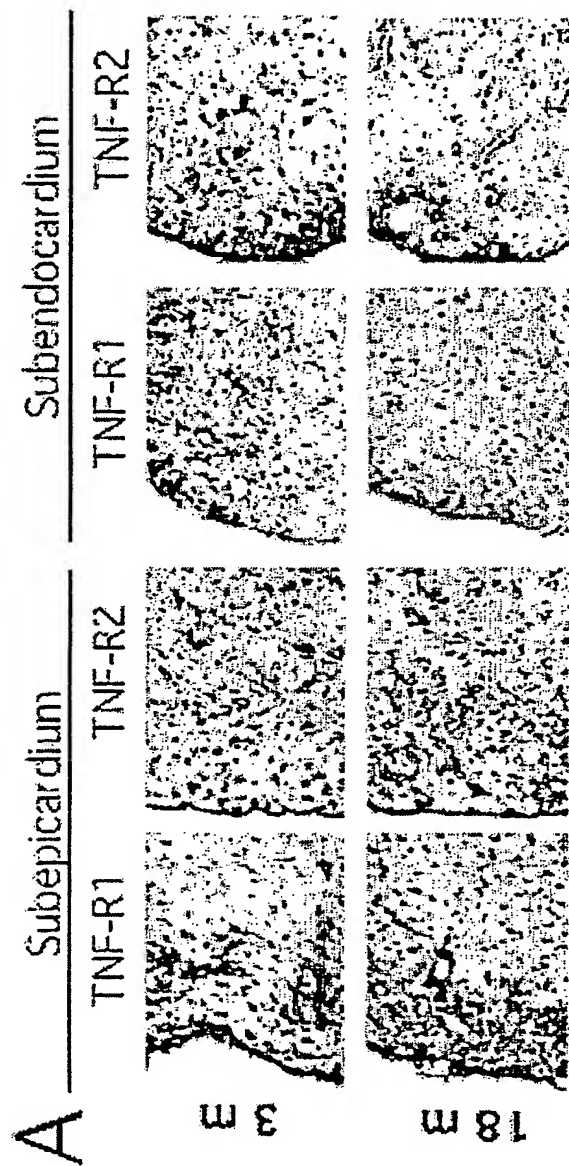


FIGURE 4B and 4C

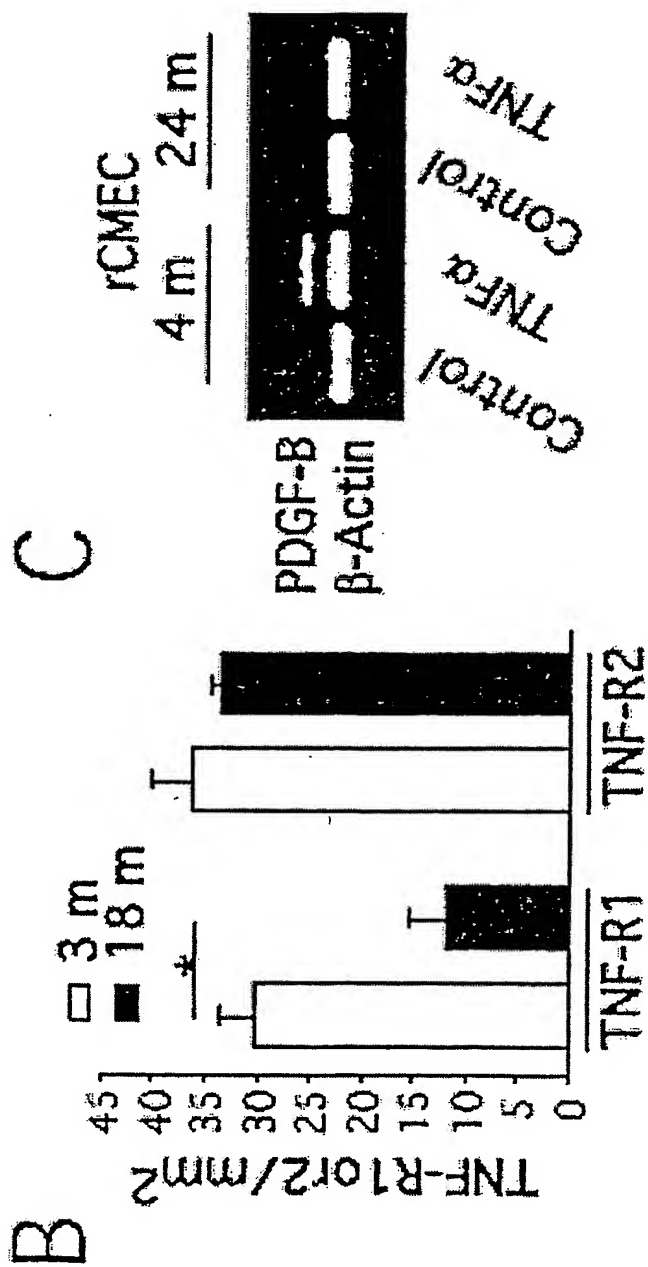


FIGURE 5

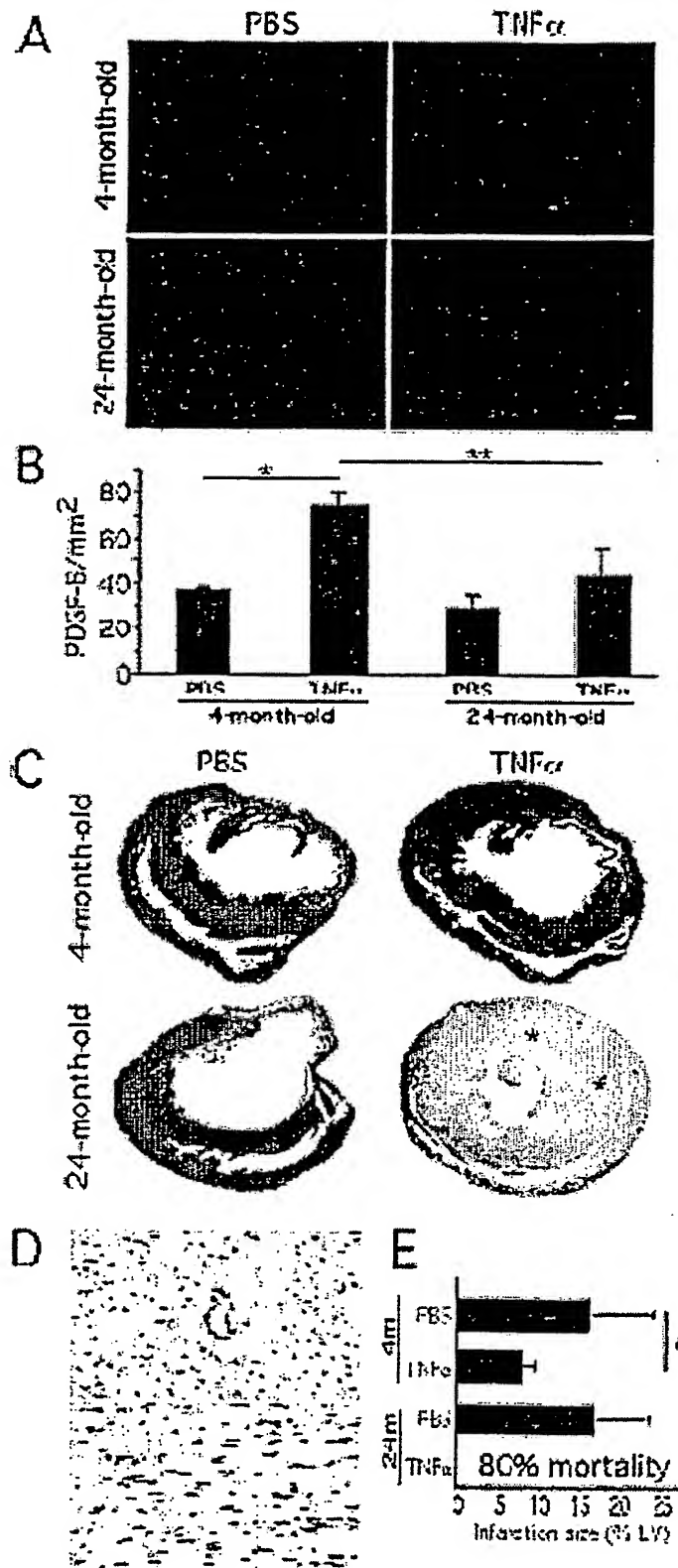


FIGURE 6

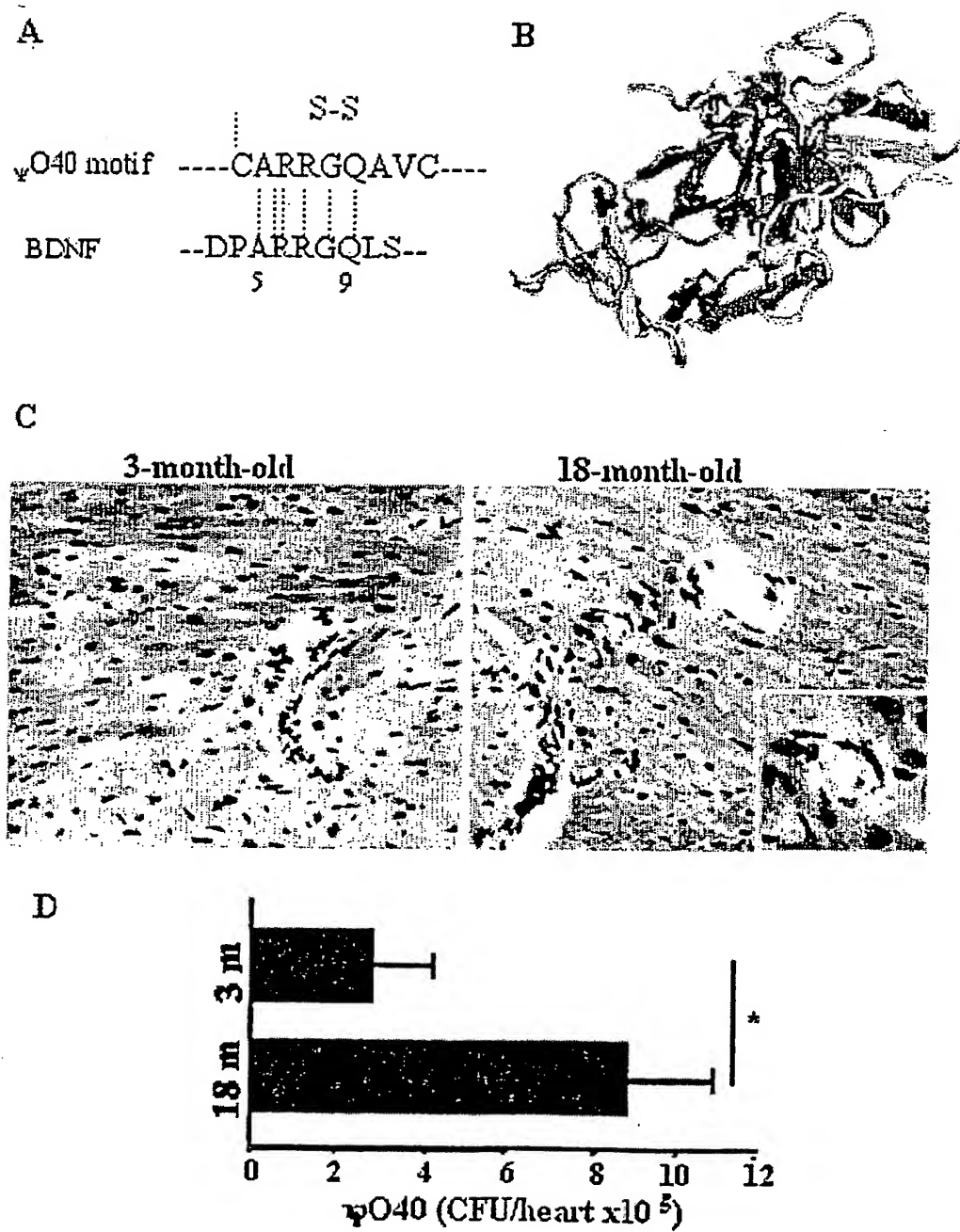


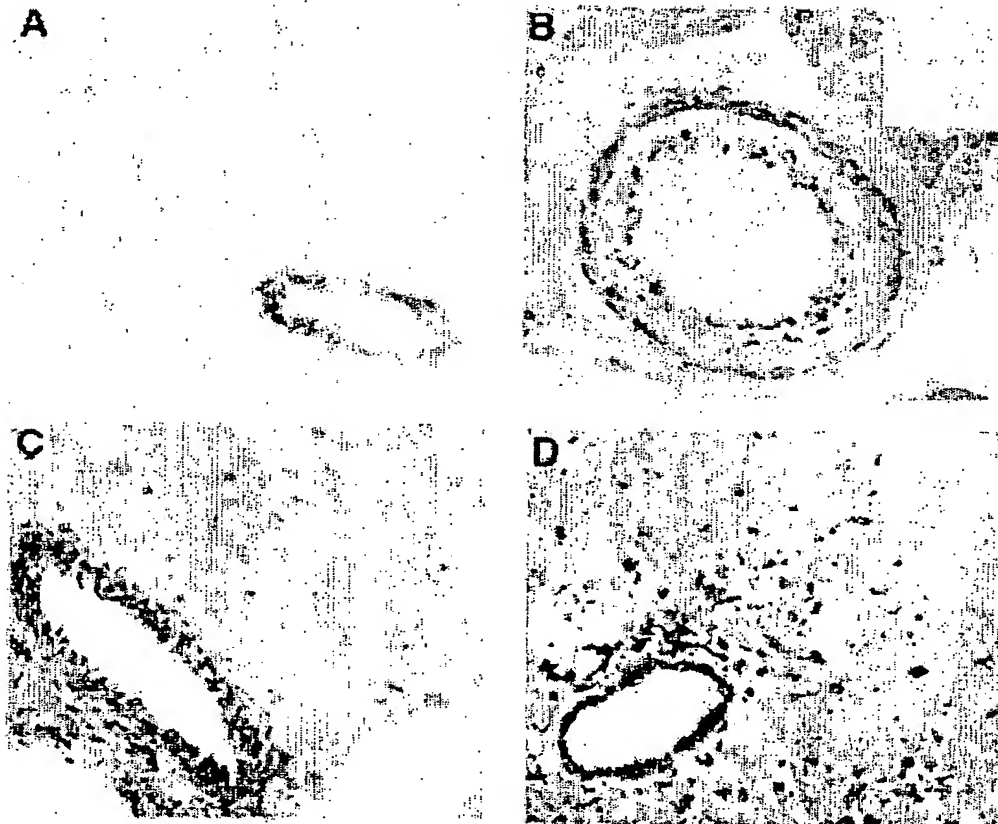
FIGURE 7**Young****Old****A****B****Trk B (extracellular)****C****D****Trk B (truncated)**

FIGURE 8

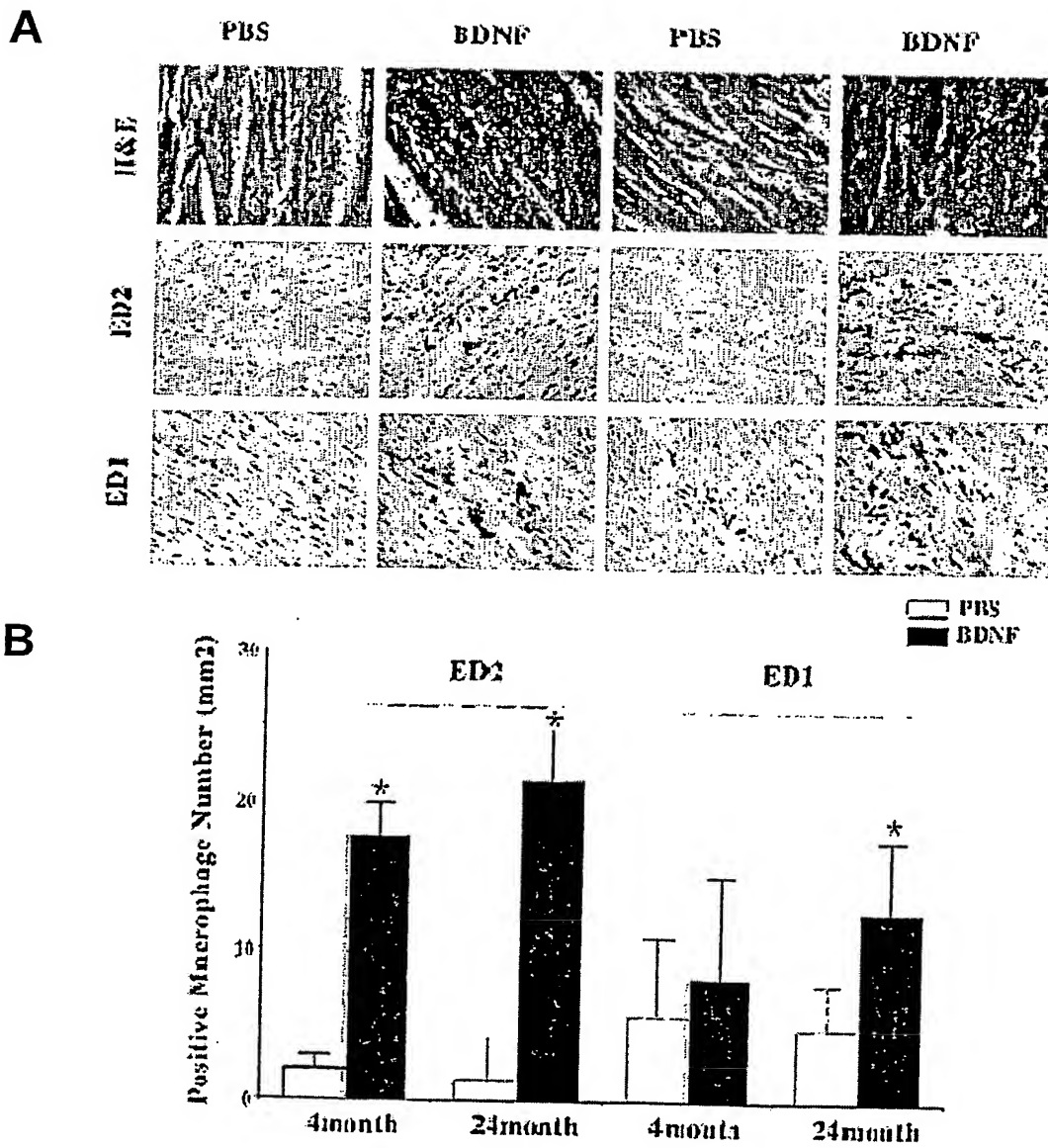


FIGURE 9A

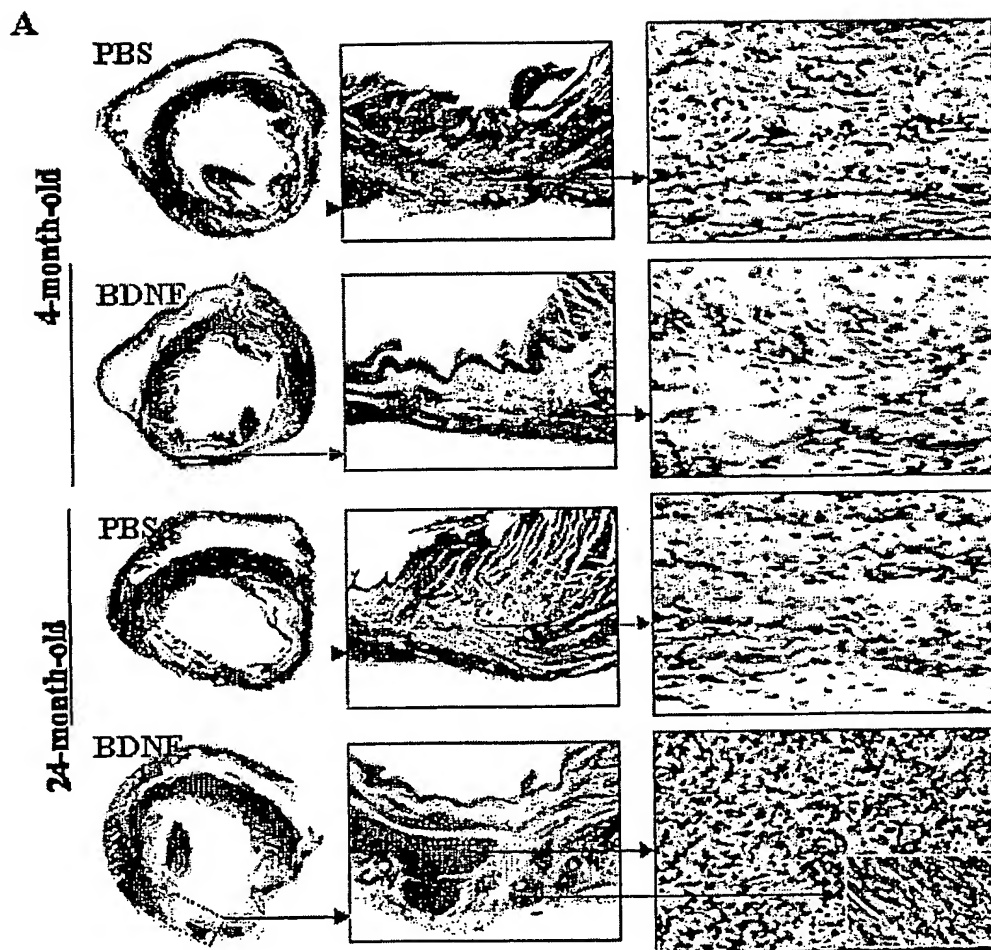


FIGURE 9B

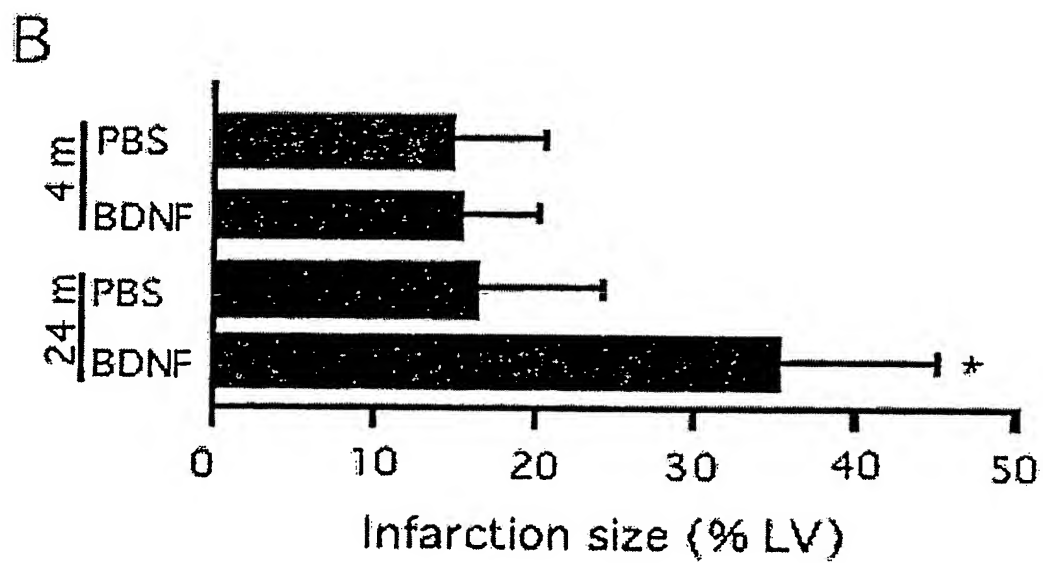
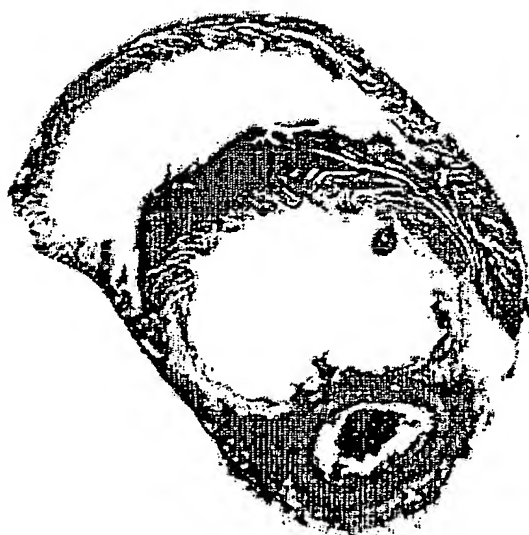


FIGURE 10

A



B

